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| APPLICATION NO.  | FILING DATE | FIRST NAMED INVENTOR | ATTORNEY DOCKET NO. | CONFIRMATION NO. |
|--|-------------|----------------------|---------------------|------------------|
| 09/993,604   | 11/14/2001  | Avi J. Ashkenazi     | P2730P1C25          | 1800             |
| 35489  | 7590        | 03/09/2004           | EXAMINER            |                  |
| HELLER EHRMAN WHITE & MCAULIFFE LLP<br>275 MIDDLEFIELD ROAD<br>MENLO PARK, CO 94025-3506 |             |                      | LANDSMAN, ROBERT S  |                  |
|  |             |                      | ART UNIT            | PAPER NUMBER     |
|  |             |                      | 1647                |                  |

DATE MAILED: 03/09/2004

Please find below and/or attached an Office communication concerning this application or proceeding.

|                              |                                      |  |  |
|------------------------------|--------------------------------------|--|--|
| <b>Office Action Summary</b> | <b>Application No.</b><br>09/993,604 | <b>Applicant(s)</b><br>GENENTECH, INC. |  |
|                              | <b>Examiner</b><br>Robert Landsman   | <b>Art Unit</b><br>1647                |  |

**-- The MAILING DATE of this communication appears on the cover sheet with the correspondence address --**

**Period for Reply**

A SHORTENED STATUTORY PERIOD FOR REPLY IS SET TO EXPIRE 3 MONTH(S) FROM THE MAILING DATE OF THIS COMMUNICATION.

- Extensions of time may be available under the provisions of 37 CFR 1.136(a). In no event, however, may a reply be timely filed after SIX (6) MONTHS from the mailing date of this communication.
- If the period for reply specified above is less than thirty (30) days, a reply within the statutory minimum of thirty (30) days will be considered timely.
- If NO period for reply is specified above, the maximum statutory period will apply and will expire SIX (6) MONTHS from the mailing date of this communication.
- Failure to reply within the set or extended period for reply will, by statute, cause the application to become ABANDONED (35 U.S.C. § 133). Any reply received by the Office later than three months after the mailing date of this communication, even if timely filed, may reduce any earned patent term adjustment. See 37 CFR 1.704(b).

**Status**

- 1) ☐ Responsive to communication(s) filed on \_\_\_\_.
- 2a) ☐ This action is **FINAL**.                      2b) ☒ This action is non-final.
- 3) ☐ Since this application is in condition for allowance except for formal matters, prosecution as to the merits is closed in accordance with the practice under *Ex parte Quayle*, 1935 C.D. 11, 453 O.G. 213.

**Disposition of Claims**

- 4) ☒ Claim(s) 119-131 is/are pending in the application.
- 4a) Of the above claim(s) \_\_\_\_ is/are withdrawn from consideration.
- 5) ☐ Claim(s) \_\_\_\_ is/are allowed.
- 6) ☒ Claim(s) 119-131 is/are rejected.
- 7) ☐ Claim(s) \_\_\_\_ is/are objected to.
- 8) ☐ Claim(s) \_\_\_\_ are subject to restriction and/or election requirement.

**Application Papers**

- 9) ☒ The specification is objected to by the Examiner.
- 10) ☐ The drawing(s) filed on \_\_\_\_ is/are: a) ☐ accepted or b) ☐ objected to by the Examiner.  
     Applicant may not request that any objection to the drawing(s) be held in abeyance. See 37 CFR 1.85(a).  
     Replacement drawing sheet(s) including the correction is required if the drawing(s) is objected to. See 37 CFR 1.121(d).
- 11) ☐ The oath or declaration is objected to by the Examiner. Note the attached Office Action or form PTO-152.

**Priority under 35 U.S.C. § 119**

- 12) ☐ Acknowledgment is made of a claim for foreign priority under 35 U.S.C. § 119(a)-(d) or (f).
- a) ☐ All    b) ☐ Some \*    c) ☐ None of:
1. ☐ Certified copies of the priority documents have been received.
2. ☐ Certified copies of the priority documents have been received in Application No. \_\_\_\_.
3. ☐ Copies of the certified copies of the priority documents have been received in this National Stage application from the International Bureau (PCT Rule 17.2(a)).

\* See the attached detailed Office action for a list of the certified copies not received.

**Attachment(s)**

- |   |  |
|---|--|
| 1) <input checked="" type="checkbox"/> Notice of References Cited (PTO-892)   | 4) <input type="checkbox"/> Interview Summary (PTO-413)<br>Paper No(s)/Mail Date. ____ |
| 2) <input type="checkbox"/> Notice of Draftsperson's Patent Drawing Review (PTO-948)  | 5) <input type="checkbox"/> Notice of Informal Patent Application (PTO-152)            |
| 3) <input checked="" type="checkbox"/> Information Disclosure Statement(s) (PTO-1449 or PTO/SB/08)<br>Paper No(s)/Mail Date <u>8/5/02</u> . | 6) <input checked="" type="checkbox"/> Other: <u>Sequence Comparison</u>               |

## **DETAILED ACTION**

### ***1. Formal Matters***

- A. The Preliminary Amendment dated 11/14/01, has been entered into the record.
- B. The Preliminary Amendment dated 9/3/02, has been entered into the record.
- C. Claims 119-131 are pending and are the subject of this Office Action.

### ***2. Priority***

Due to the excessive number of applications from which the present application claims benefit, priority cannot be determined. However, the Examiner has concluded that the subject matter defined in this application is not supported by any of the applications in the chain of priority because the presently claimed subject matter is not supported by a specific, substantial or well-established utility, nor, for this reason, is it enabled. Accordingly, the subject matter defined in claims 119-131 has an effective filing date of 11/14/01, which is the filing date of the present application.

Should the applicant disagree with the examiner's factual determination above, it is incumbent upon the applicant to provide the serial number and specific page number(s) of any parent application filed prior to 11/14/01 which specifically supports the particular claim limitation for each and every claim limitation in all the pending claims which applicant considers to have been in possession of and fully enabled for prior to 11/14/01.

### ***3. Information Disclosure Statement***

- A. References A1 and A2 on the IDS filed 5/24/02 have been lined through since they are not in proper format, including author and date of deposit.

### ***4. Specification***

- A. Though none could be found, due to the length of the specification, Applicants are reminded that embedded hyperlink and/or other form of browser-executable code are not permitted in the specification. See MPEP § 608.01.

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B. The title of the invention is not descriptive. A new title is required that is clearly indicative of the invention to which the claims are directed. The title recites polypeptides and polynucleotides whereas the claims are drawn to polypeptides.

### **5. Claim Objections**

A. The syntax of claims 119-131 could be improved by replacing the phrase “shown in Figure 233 (SEQ ID NO:326)” with “of SEQ ID NO:326.”

### **6. Claim Rejections - 35 USC § 101**

35 U.S.C. 101 reads as follows:

Whoever invents or discovers any new and useful process, machine, manufacture, or composition of matter, or any new and useful improvement thereof, may obtain a patent therefor, subject to the conditions and requirements of this title.

A. Claims 119-131 are rejected under 35 U.S.C. 101 because the claimed invention is not supported by a specific, substantial and credible asserted utility or a well established utility. These claims are directed to polypeptides having various sequence homology to SEQ ID NO:326. However, the invention encompassed by these claims has no apparent or disclosed patentable utility. This rejection is consistent with the current utility guidelines, published 1/5/01, 66 FR 1092. The instant application has provided a description of an isolated protein. However, the instant application does not disclose a specific and substantial biological role of this protein or its significance.

However, it is clear from the instant specification that the claimed protein is what is termed an “orphan receptor” in the art. The instant application does not disclose the biological role of the claimed protein or its significance. Applicants disclose in the specification that the receptor is a secreted protein. However, this fact, alone, is insufficient to confer utility to the protein of the present invention. Therefore, the instant claims are drawn to a polynucleotide encoding a protein which has a yet undetermined function or biological significance. There is no actual and specific significance which can be attributed to said protein identified in the specification. For this reason, the instant invention is incomplete. In the absence of a knowledge of the natural ligands or biological significance of this protein, there is no immediately obvious patentable use for it. To employ a protein of the instant invention in the identification of substances which bind to and/or mediate activity of the said receptor is clearly to use it as the object of further research which has been determined by the courts to be a non-patentable utility. Since the instant specification does not disclose a “real-world” use for said protein then the claimed invention is incomplete and, therefore, does not meet the requirements of 35 U.S.C. 101 as being useful.

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**Furthermore, since the protein of the invention is not supported by a specific and substantial asserted utility or a well established utility, the encoding polynucleotides and chimeric proteins also lack utility.**

***7. Claim Rejections - 35 USC § 112, first paragraph - enablement***

The following is a quotation of the first paragraph of 35 U.S.C. 112:

The specification shall contain a written description of the invention, and of the manner and process of making and using it, in such full, clear, concise, and exact terms as to enable any person skilled in the art to which it pertains, or with which it is most nearly connected, to make and use the same and shall set forth the best mode contemplated by the inventor of carrying out his invention.

A. Claims 119-131 are rejected under 35 U.S.C. 112, first paragraph, as failing to adequately teach how to use the instant invention. Specifically, since the claimed invention is not supported by a specific, substantial and credible asserted utility or a well established utility for the reasons set forth above, one skilled in the art clearly would not know how to use the claimed invention.

B. Claims 119-131 are rejected under 35 U.S.C. 112, first paragraph, as containing subject matter which was not described in the specification in such a way as to enable one skilled in the art to which it pertains, or with which it is most nearly connected, to make and/or use the invention.

The deposit of the biological material is considered necessary for the enablement of the current invention (see MPEP Chapter 2400 and 37 C.F.R. §§ 1.801-1.809). Elements required for practicing a claimed invention must be known and readily available to the public or obtainable by a repeatable method set forth in the specification. If a deposit (203129) is made under the terms of the Budapest Treaty on the International Recognition of the Deposit of Microorganisms for the Purposes of Patent Procedure (e.g. see 961 OG 21, 1977), and Applicants, their assignee or their agent needs to provide a declaration containing the following:

1. the current address of the ATCC.
2. a declaration, or statement over attorney's signature stating that all restrictions imposed by the depositor on the availability to the public of the deposited biological material be irrevocably removed upon the granting of the patent (see MPEP Chapter 2410.01 and 37 C.F.R. § 1.808).

C. Furthermore, even if the claims possessed utility under 35 USC 101, claims 119-131 would still be rejected under 35 USC 112, first paragraph, because the specification, while then being enabling for SEQ ID NO:325 and 326, does not reasonably provide enablement for polypeptides having at least 80%, 85%, 90%, 95% or 99% sequence identity to SEQ ID NO:326, to the protein encoded by ATCC No.

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203129, for the extracellular domain thereof, or for fusion proteins. The specification does not enable any person skilled in the art to which it pertains, or with which it is most nearly connected, to make the invention commensurate in scope with these claims. There is no functional limitation in the claims. The claims encompass an unreasonable number of inoperative polypeptides, or polynucleotides which encode these polypeptides, which the skilled artisan would not know how to use.

There are no working examples of polynucleotides or polypeptides less than 100% identical to SEQ ID NO:325 or 326, or the mature form thereof (i.e. lacking its signal peptide). The skilled artisan would not know how to use non-identical polypeptides on the basis of teachings in the prior art or specification unless they possessed a specific function disclosed in the instant specification, in which there is none. While the specification generally describes homologous proteins, Applicants still have not taught to which family of proteins the protein of the present invention belongs. The specification does not provide guidance for using polynucleotides encoding polypeptides related to (i.e., 80%-99% identity) but not identical to SEQ ID NO:325 or 326 which do not have any specific, known function. The claims are broad because they do not require the claimed polypeptide to be identical to the disclosed sequence and because the claims have no functional limitation.

For these reasons, which include the complexity and unpredictability of the nature of the invention and art in terms of the diversity of proteases and lack of knowledge about function(s) of encompassed polypeptides structurally related to SEQ ID NO:325, the lack of direction or guidance for using polypeptides that are not identical to SEQ ID NO:326, and the breadth of the claims for structure without function, it would require undue experimentation to use the invention commensurate in scope with the claims.

#### ***8. Claim Rejections - 35 USC § 112, first paragraph – written description***

A. Claims 119-131 are rejected under 35 U.S.C. 112, first paragraph, as containing subject matter which was not described in the specification in such a way as to reasonably convey to one skilled in the relevant art that the inventor(s), at the time the application was filed, had possession of the claimed invention.

The claims are drawn to polypeptides having at least 80%, 85%, 90%, 95% or 99% sequence identity with SEQ ID NO:326, and fusion proteins thereof. The claims do not require that the polypeptide of the present invention possess any particular biological activity, nor any particular conserved structure, or other disclosed distinguishing feature. Thus, the claims are drawn to a genus of polypeptides that is defined only by sequence identity.

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To provide adequate written description and evidence of possession of a claimed genus, the specification must provide sufficient distinguishing identifying characteristics of the genus. The factors to be considered include disclosure of complete or partial structure, physical and/or chemical properties, functional characteristics, structure/function correlation, methods of making the claimed product, or any combination thereof. In this case, the only factor present in the claim is a partial structure in the form of a recitation of percent identity. There is not even identification of any particular portion of the structure that must be conserved. Accordingly, in the absence of sufficient recitation of distinguishing identifying characteristics, the specification does not provide adequate written description of the claimed genus.

*Vas-Cath Inc. v. Mahurkar*, 19USPQ2d 1111, clearly states “applicant must convey with reasonable clarity to those skilled in the art that, as of the filing date sought, he or she was in possession of the invention. The invention is, for purposes of the ‘written description’ inquiry, *whatever is now claimed*.” (See page 1117.) The specification does not “clearly allow persons of ordinary skill in the art to recognize that [he or she] invented what is claimed.” (See *Vas-Cath* at page 1116). As discussed above, the skilled artisan cannot envision the detailed chemical structure of the encompassed genus of polypeptides, and therefore conception is not achieved until reduction to practice has occurred, regardless of the complexity or simplicity of the method of isolation. Adequate written description requires more than a mere statement that it is part of the invention and reference to a potential method of isolating it. The compound itself is required. See *Fiers v. Revel*, 25 USPQ2d 1601 at 1606 (CAFC 1993) and *Amgen Inc. v. Chugai Pharmaceutical Co. Ltd.*, 18 USPQ2d 1016.

One cannot describe what one has not conceived. See *Fiddes v. Baird*, 30 USPQ2d 1481 at 1483. In *Fiddes*, claims directed to mammalian FGF’s were found to be unpatentable due to lack of written description for that broad class. The specification provided only the bovine sequence.

Therefore, only isolated polypeptides comprising the amino acid sequence set forth in SEQ ID NO:326, or encoded by SEQ ID NO:325, but not the full breadth of the claims, meets the written description provision of 35 U.S.C. §112, first paragraph. Applicant is reminded that *Vas-Cath* makes clear that the written description provision of 35 U.S.C. §112 is severable from its enablement provision (see page 1115).

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**9. Claim Rejections - 35 USC § 112, second paragraph**

The following is a quotation of the second paragraph of 35 U.S.C. 112:

The specification shall conclude with one or more claims particularly pointing out and distinctly claiming the subject matter which the applicant regards as his invention.

Claims 119-131 are rejected under 35 U.S.C. 112, second paragraph, as being indefinite for failing to particularly point out and distinctly claim the subject matter which applicant regards as the invention.

A. Claims 119-131 are vague and indefinite since it is not clear whether or not the protein of the present invention is a soluble protein (e.g protease), nor is it disclosed as being expressed on a cell surface. Accordingly, the limitation that the claimed protein comprises an “extracellular domain” is indefinite, as the art does not recognize soluble proteins as having such domains. Further, if the protein had an extracellular domain, the recitation of “the extracellular domain”...”lacking its associated signal sequence” is indefinite as a signal sequence is not generally considered to be part of an extracellular domain, as signal sequences are cleaved from said domains in the process of secretion from the cell.

**10. Claim Rejections - 35 USC § 102**

The following is a quotation of the appropriate paragraphs of 35 U.S.C. 102 that form the basis for the rejections under this section made in this Office action:

A person shall be entitled to a patent unless –

(b) the invention was patented or described in a printed publication in this or a foreign country or in public use or on sale in this country, more than one year prior to the date of application for patent in the United States.

A. Claims 119-131 are rejected under 35 U.S.C. 102(b) as being anticipated by Baker et al. (WO 99/63088). The claims recite an isolated polypeptide at least 80% identical to SEQ ID NO:326 as well as polynucleotides encoding this protein, extracellular domains and chimeric polypeptides. Baker et al. teach a protein which is 100% identical to SEQ ID NO:326 of the present invention (Sequence Comparison). This protein would encompass all of the claimed variants of that of the present invention. Baker also teach chimeric peptides (page 350, line 15).

**11. Other Pertinent Art**

A. Tang et al. (WO 01/53312) teach a protein which is 54.7% identical to that of SEQ ID NO:326 of the present invention (Sequence Comparison A).



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B. Wiemann et al. (Genome Research) teach a protein which is 67.1% identical to that of SEQ ID NO:326 of the present invention (Sequence Comparison B).

**12. Conclusion**

A. No claim is allowable.

***Advisory information***

Any inquiry concerning this communication or earlier communications from the examiner should be directed to Robert Landsman whose telephone number is (703) 306-3407. The examiner can normally be reached on Monday - Friday from 8:00 AM to 5:00 PM (Eastern time) and alternate Fridays from 8:00 AM to 5:00 PM (Eastern time).

If attempts to reach the examiner by telephone are unsuccessful, the Examiner's supervisor, Gary Kunz, can be reached on (703) 308-4623.

Official papers filed by fax should be directed to (703) 308-4242. Fax draft or informal communications with the examiner should be directed to (703) 308-0294.

Any inquiry of a general nature or relating to the status of this application or proceeding should be directed to the Group receptionist whose telephone number is (703) 308-0196.

Robert Landsman, Ph.D.  
Patent Examiner  
Group 1600  
March 04, 2004

  
ROBERT LANDSMAN  
PATENT EXAMINER

# SEQ ID NO: 326

# Sequence Comparison

ID AAY66729 standard; protein; 775 AA.  
 XX  
 AC AAY66729;  
 XX  
 DT 05-APR-2000 (first entry)  
 XX  
 DE Membrane-bound protein PRO1281.  
 XX  
 KW Membrane-bound polypeptide; PRO polypeptide; LDL receptor; TIE ligand;  
 KW pharmaceutical; receptor immunoadhesin; gene mapping.  
 XX  
 OS Homo sapiens.  
 XX  
 PN WO9963088-A2.  
 XX  
 PD 09-DEC-1999.  
 XX  
 PF 02-JUN-1999; 99WO-US12252.  
 XX  
 PR 02-JUN-1998; 98US-0087607.  
 XX  
 PA (GETH ) GENENTECH INC.  
 XX  
 PI Baker K, Chen J, Goddard A, Gurney AL, Smith V, Watanabe CK;  
 PI Wood WI, Yuan J;  
 XX  
 DR WPI; 2000-072883/06.  
 DR N-PSDB; AAZ65074.  
 XX  
 PT Membrane-bound proteins and related nucleotide sequences -  
 XX  
 PS claim 12; Fig 233; 822pp; English.  
 XX  
 CC The invention provides membrane-bound PRO polypeptides and  
 CC polynucleotides encoding them. The PRO sequences of the invention were  
 CC identified based on extracellular domain homology screening. The PRO  
 CC sequences have homology with proteins including LDL receptors, TIE  
 CC ligands and various enzymes. The membrane-bound proteins and receptor  
 CC molecules are useful as pharmaceutical and diagnostic agents. Receptor  
 CC immunoadhesins, for instance, can be used as therapeutic agents to block  
 CC receptor-ligand interactions. The membrane-bound proteins can also be  
 CC employed for screening of potential peptide or small molecule inhibitors  
 CC of the relevant receptor/ligand interaction. The PRO encoding sequences  
 CC are useful as hybridization probes, in chromosome and gene mapping and in  
 CC the generation of antisense RNA and DNA. PRO nucleic acid sequences  
 CC will also be useful for the preparation of PRO polypeptides, especially  
 CC by recombinant techniques.  
 XX  
 SQ Sequence 775 AA;

Query Match 100.0%; Score 4074; DB 21; Length 775;  
 Best Local Similarity 100.0%; Pred. No. 0;  
 Matches 775; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 MRASLLSVLRPAGPVAVGISLGFTLSLLSVTWVEEPCGPGPPQPGDSELPPRGNTNAAR 60  
 ||||||||||||||||||||||||||||||||||||||||||||||||||||||||  
 Db 1 MRASLLSVLRPAGPVAVGISLGFTLSLLSVTWVEEPCGPGPPQPGDSELPPRGNTNAAR 60  
 QY 61 RPNSVQPGAEREKPGAGEGAGENWEPRVLPYHPAQPGAQAAKAVRTRYISTELGIRQLL 120  
 ||||||||||||||||||||||||||||||||||||||||||||||||||||||||  
 Db 61 RPNSVQPGAEREKPGAGEGAGENWEPRVLPYHPAQPGAQAAKAVRTRYISTELGIRQLL 120

|    |     |   |     |
|----|-----|---|-----|
| Qy | 121 | VAVLTSQTTLPTLGVAVNRTLGHRLERVVFLTGARGRRAPPGMAVVTLGEERPIGHLHLA  | 180 |
|    |     |   |     |
| Db | 121 | VAVLTSQTTLPTLGVAVNRTLGHRLERVVFLTGARGRRAPPGMAVVTLGEERPIGHLHLA  | 180 |
| Qy | 181 | LRHLEQHGDDFDWFFLVPDTTYTEAHGLARLTGHLSLASAAHLYLGRPQDFIGGEPTPG   | 240 |
|    |     |   |     |
| Db | 181 | LRHLEQHGDDFDWFFLVPDTTYTEAHGLARLTGHLSLASAAHLYLGRPQDFIGGEPTPG   | 240 |
| Qy | 241 | RYCHGGFGVLLSRMLLQQLRPHLEGCRNDIVSARPDEWLGRCILDATGVGCTGDHEGVHY  | 300 |
|    |     |   |     |
| Db | 241 | RYCHGGFGVLLSRMLLQQLRPHLEGCRNDIVSARPDEWLGRCILDATGVGCTGDHEGVHY  | 300 |
| Qy | 301 | SHLELSPGEPVQEGDPHFRSALTAHPVRDPVHMYQLHKAFARAEELERTYQEIQELQWEIQ | 360 |
|    |     |   |     |
| Db | 301 | SHLELSPGEPVQEGDPHFRSALTAHPVRDPVHMYQLHKAFARAEELERTYQEIQELQWEIQ | 360 |
| Qy | 361 | NTSHLAVDGDRAAAWPVGIPAPSRPASRFEVLRWDYFTEQHAFSCADGSPRCPLRGADRA  | 420 |
|    |     |   |     |
| Db | 361 | NTSHLAVDGDRAAAWPVGIPAPSRPASRFEVLRWDYFTEQHAFSCADGSPRCPLRGADRA  | 420 |
| Qy | 421 | DVADVLGTALEELNRRYHPALRLQKQQLVNGYRRFDPARGMEYTLDLQLEALTPQGGRRP  | 480 |
|    |     |   |     |
| Db | 421 | DVADVLGTALEELNRRYHPALRLQKQQLVNGYRRFDPARGMEYTLDLQLEALTPQGGRRP  | 480 |
| Qy | 481 | LTRRVQLLRPLSRVEILPVPYVTEASRLTVLLPLAAAERDLAPGFLEAFATAALEPGDAA  | 540 |
|    |     |   |     |
| Db | 481 | LTRRVQLLRPLSRVEILPVPYVTEASRLTVLLPLAAAERDLAPGFLEAFATAALEPGDAA  | 540 |
| Qy | 541 | AALTLLLLYEPRQAQRVAHADVFAPVKAHVAELERRFPGARVPWLSVQTAAPSPLRLMDL  | 600 |
|    |     |   |     |
| Db | 541 | AALTLLLLYEPRQAQRVAHADVFAPVKAHVAELERRFPGARVPWLSVQTAAPSPLRLMDL  | 600 |
| Qy | 601 | LSKKHPLDTLFLLAGPDTVLTDPFLNRCRMHAISGWQAFFPMHFQAFHPGVAPPQGP     | 660 |
|    |     |   |     |
| Db | 601 | LSKKHPLDTLFLLAGPDTVLTDPFLNRCRMHAISGWQAFFPMHFQAFHPGVAPPQGP     | 660 |
| Qy | 661 | ELGRDTGRFDRQAASEACFYNSDYVAARGRLAAASEQEEELLESLDVYELFLHFSSHLVL  | 720 |
|    |     |   |     |
| Db | 661 | ELGRDTGRFDRQAASEACFYNSDYVAARGRLAAASEQEEELLESLDVYELFLHFSSHLVL  | 720 |
| Qy | 721 | RAVEPALLQRYRAQTCSARLSEDLYHRCLQSVLEGLGSRTQLAMLLFEQE QGNST      | 775 |
|    |     |   |     |
| Db | 721 | RAVEPALLQRYRAQTCSARLSEDLYHRCLQSVLEGLGSRTQLAMLLFEQE QGNST      | 775 |

# Sequence Comparison A

ID AAM39781 standard; Protein; 772 AA.  
 XX  
 AC AAM39781;  
 XX  
 DT 22-OCT-2001 (first entry)  
 XX  
 DE Human polypeptide SEQ ID NO 2926.  
 XX  
 KW Human; nootropic; immunosuppressant; cytostatic; gene therapy; cancer;  
 KW peripheral nervous system; neuropathy; central nervous system; CNS;  
 KW Alzheimer's; Parkinson's disease; Huntington's disease; haemostatic;  
 KW amyotrophic lateral sclerosis; Shy-Drager Syndrome; chemotactic;  
 KW chemokinetic; thrombolytic; drug screening; arthritis; inflammation;  
 KW leukaemia.  
 XX  
 OS Homo sapiens.  
 XX  
 PN WO200153312-A1.  
 XX  
 PD 26-JUL-2001.  
 XX  
 PF 26-DEC-2000; 2000WO-US34263.  
 XX  
 PR 21-JAN-2000; 2000US-0488725.  
 PR 25-APR-2000; 2000US-0552317.  
 PR 09-JUL-2000; 2000US-0598042.  
 PR 19-JUL-2000; 2000US-0620312.  
 PR 03-AUG-2000; 2000US-0653450.  
 PR 14-SEP-2000; 2000US-0662191.  
 PR 19-OCT-2000; 2000US-0693036.  
 PR 29-NOV-2000; 2000US-0727344.  
 XX  
 PA (HYSE-) HYSEQ INC.  
 XX  
 PI Tang YT, Liu C, Asundi V, Chen R, Ma Y, Qian XB, Ren F, Wang D;  
 PI Wang J, Wang Z, Wehrman T, Xu C, Xue AJ, Yang Y, Zhang J;  
 PI Zhao QA, Zhou P, Goodrich R, Drmanac RT;  
 XX  
 DR WPI; 2001-442253/47.  
 DR N-PSDB; AAI58937.  
 XX  
 PT Novel nucleic acids and polypeptides, useful for treating disorders  
 PT such as central nervous system injuries -  
 XX  
 PS Example 4; SEQ ID NO 2926; 10078pp; English.  
 XX  
 CC The invention relates to human nucleic acids (AAI57798-AAI61369) and  
 CC the encoded polypeptides (AAM38642-AAM42213) with nootropic,  
 CC immunosuppressant and cytostatic activity. The polynucleotides are useful  
 CC in gene therapy. A composition containing a polypeptide or polynucleotide  
 CC of the invention may be used to treat diseases of the peripheral nervous  
 CC system, such as peripheral nervous injuries, peripheral neuropathy and  
 CC localised neuropathies and central nervous system diseases, such as  
 CC Alzheimer's, Parkinson's disease, Huntington's disease, amyotrophic  
 CC lateral sclerosis, and Shy-Drager Syndrome. Other uses include the  
 CC utilisation of the activities such as: Immune system suppression,  
 CC Activin/inhibin activity, chemotactic/chemokinetic activity, haemostatic  
 CC and thrombolytic activity, cancer diagnosis and therapy, drug screening,  
 CC assays for receptor activity, arthritis and inflammation, leukaemias and  
 CC C.N.S disorders.  
 CC Note: The sequence data for this patent did not form part of the printed  
 CC specification.  
 XX

SQ Sequence 772 AA;

Query Match 54.7%; Score 2227.5; DB 22; Length 772;  
Best Local Similarity 56.6%; Pred. No. 1.2e-180;  
Matches 455; Conservative 101; Mismatches 187; Indels 61; Gaps 14;

```
Qy      1 MRASLLLSVLRPAGPVAVGISLGFTLSLLSVTWV---EEPC----GPGPPQPGDSELP 51
      || | ||::||| | : :||| :||| | :| : | | | |
Db      1 MRLSSLLALLRPALPLILGLSLGCSLLRVSWIQGEGEDPCVEAVGERGGPQNPSR-- 58

Qy     52 PRGNTNAARRPNSVQPGAEREKPGAGEGAGENWEPRVLPYHPAQPGQAACKAVRTRYIST 111
      || | | | | | | | | | | | | | | | | | | | | | |
Db     59 -----ARLDQS-----DEDFKPRIVPYR-RDPNKPYPKVLTRYIQT 94

Qy    112 ELGIRQRLLVAVLTSQTTLPTLGVAVNRTLGHRLERVVFLTGARGRRAPPGMAVVTLGEE 171
      || | | :||| ||||| : || | | ||||| : | | : : || | | | | | | |
Db    95 ELGSRELLVAVLTSRATLSTLAVAVNRTVAHHFPRLLYFTGQRCARAPAGMQVVSNGDE 154

Qy    172 RPIGHLHLALRHLLLEQHGDGDFWFFLVPDTTYTEAHGLARLTGHLSLASAAHYLGRPD 231
      || : | | | | | | | | | | | | | | | | | | | | | | | |
Db    155 RPAWLMSETLRHLHTHFGADYDWFIMQDDTYVQAPRLAALAGHLSINQ--DLYLGRAEE 212

Qy    232 FIG-GEPTPGRYCHGGFGVLLSRMLLQQLRPHLEGCRNDIVSARPDEWLGRCLDATGVG 290
      || | | | | | | | | | | | | | | | | | | | | | | | |
Db    213 FIGAGE--QARYCHGGFGYLLSRSLLLRLRPHLDGCRGDILSARPDEWLGRCLIDSLGVG 270

Qy    291 CTGDHEGVHYSHLELSPG-EPVQEGDPHFRSALTAHPVRDPVHMYQLHKAFARAELETTY 349
      | : | | | | | | | | : : | | | | | | | | | | : | | | | |
Db    271 CVSQHQQQYRSFELAKNRDPEKEGSSAFLSAFVHPVSEGTLMYRLHKRFSALELERAY 330

Qy    350 QEIQELQWEIQNTSHLAVDGDRAAAWPVGI PAPS RPASRFEVLRWDYFTEQHAFSCADGS 409
      || : || : | : | : : : ||| : || | | | | | | | | | | | :
Db    331 SEIEQLQAQIRNLTVLTPGEAGLSWPVGLPAPFTPHSRFEVLGWDYFTEQHTFSCADGA 390

Qy    410 PRCPLRGADRADVADVLGTALEELNRRYPALRLQKQLVNGYRRFDPARGMEYTLDLQL 469
      | : || : || | | | | | | | | | | | | | | | | | | | | | |
Db    391 PKCPLQGASRADVGDALETALEQLNRRYQPRLRFPQKQRLNGYRRFDPARGMEYTLDLLL 450

Qy    470 EALTPQGGRRPLTRRVQLLRPLSRVEILPVPYVTEASRLTVLLPLAAERDLAPGFLEAF 529
      | : | : | | | | | | | | | | | | | | | | : : || | | | | |
Db    451 ECVTQRGHRRALARRVSLRPLSRVEILPMPYVTEATRVQLVPLLVAAAAAPAFLEAF 510

Qy    530 ATAALPGDAAAALTLLLLYEPRQAQRVAHADVFAPVKAHVAELERRFPGARVPWLSVQT 589
      | | | : | | | | : | | : | | | | | | | | | | : | | : |
Db    511 AANVLEPRE-HALLTLLLVYGPREGGRGA-PDPFLGVKAAAELERRYPGTRLAWLAVRA 568

Qy    590 AAPSPRLRLMDLLSKKHPLDTLFLLAGPDTVLTDPFLNRCRMHAISGWQAFFPMHFQAFHP 649
      || | : ||| : : ||| : || | | | | | | | | | | | | | | | |
Db    569 EAPSQVRLMDVVSKKHVPDVLFFLTWTRPGPEVLNRCRMNAISGWQAFFPVHFQEFNP 628

Qy    650 GVAPPQG-PGPPPELGRDT-----GRFDRQAASEACFYNSDYVAARGRLAA 693
      : : | : | | | | | | | | | | | | | | : : | | : | | | |
Db    629 ALSPQRSPPGPPGAGDPFPPSPGADPSRGAPIGGRFDRQASAEGCFYNADYLAARLAG 688

Qy    694 --ASEQEEELLES LDVYELFLHFSSLHVLRAVEPALLQRYRAQTCSARLSEDLYHRC LQS 751
      | : || | | | : | : || | | | : || | | | : : : | | | | : || |
Db    689 ELAQEEEEALEGLEVMDFLRFSGHLHFRAVEPGLVQKFSLRDCSPRLSEELYHRCRLS 748

Qy    752 VLEGLGSRTQLAMLLFEQE QGNST 775
      |||| | | | | | | | | |
Db    749 NLEGLGGRAQLAMALFEQE QANST 772
```

# Sequence Comparison B

ID Q9H0F8 PRELIMINARY; PRT; 522 AA.  
AC Q9H0F8;  
DT 01-MAR-2001 (TrEMBLrel. 16, Created)  
DT 01-MAR-2001 (TrEMBLrel. 16, Last sequence update)  
DT 01-OCT-2002 (TrEMBLrel. 22, Last annotation update)  
DE Hypothetical protein.  
GN DKFZP434E0423.  
OS Homo sapiens (Human).  
OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;  
OC Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.  
OX NCBI\_TaxID=9606;  
RN [1]  
RP SEQUENCE FROM N.A.  
RC TISSUE=Testis;  
RX MEDLINE=21154917; PubMed=11230166;  
RA Wiemann S., Weil B., Wellenreuther R., Gassenhuber J., Glassl S.,  
RA Ansorge W., Boecher M., Bloecker H., Bauersachs S., Blum H.,  
RA Lauber J., Duesterhoeft A., Beyer A., Koehrer K., Strack N.,  
RA Mewes H.W., Ottenwaelde B., Obermaier B., Tampe J., Heubner D.,  
RA Wambutt R., Korn B., Klein M., Poustka A.;  
RT "Towards a Catalog of Human Genes and Proteins: Sequencing and  
RT Analysis of 500 Novel Complete Protein Coding Human cDNAs.";  
RL Genome Res. 11:422-435(2001).  
RN [2]  
RP SEQUENCE FROM N.A.  
RC TISSUE=Lymph;  
RA Strausberg R.;  
RL Submitted (AUG-2001) to the EMBL/GenBank/DDBJ databases.  
RN [3]  
RP SEQUENCE FROM N.A.  
RC TISSUE=Brain;  
RA Strausberg R.;  
RL Submitted (JAN-2002) to the EMBL/GenBank/DDBJ databases.  
DR EMBL; AL136814; CAB66748.1; -.  
DR EMBL; BC013369; AAH13369.1; -.  
DR EMBL; BC021223; AAH21223.1; -.  
KW Hypothetical protein.  
SQ SEQUENCE 522 AA; 58355 MW; 87501CE0A043AE3B CRC64;

Query Match 67.1%; Score 2732; DB 4; Length 522;  
Best Local Similarity 99.8%; Pred. No. 3e-184;  
Matches 521; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

|    |     |   |     |
|----|-----|---|-----|
| Qy | 254 | MLLQQLRPHLEGCRNDIVSARPDEWLGRCLDATGVGCTGDHEGVHYSHLELSPGEPVQE   | 313 |
| Db | 1   | MLLQQLRPHLEGCRNDIVSARPDEWLGRCLDATGVGCTGDHEGVHYSHLELSPGEPVQE   | 60  |
| Qy | 314 | GDPHFRSALTAHPVRDPVHMYQLHKAFARAELEPTYQEIQELQWEIQNTSHLAVDGDRAA  | 373 |
| Db | 61  | GDPHFRSALTAHPVRDPVHMYQLHKAFARAELEPTYQEIQELQWEIQNTSHLAVDGDRAA  | 120 |
| Qy | 374 | AWPVGIPAPSRPASRFEVLRWDYFTEQHAFSCADGSPRCPLRGADRADVADVLTGTALEEL | 433 |
| Db | 121 | AWPVGIPAPSRPASRFEVLRWDYFTEQHAFSCADGSPRCPLRGADRADVADVLTGTALEEL | 180 |
| Qy | 434 | NRRYHPALRLQKQQLVNGYRRFDPARGMEYTLDLQLEALTPQGRRPLTRRVQLLRPLSR   | 493 |
| Db | 181 | NRRYHPALRLQKQQLVNGYRRFDPARGMEYTLDLQLEALTPQGRRPLTRRVQLLRPLSR   | 240 |
| Qy | 494 | VEILPVPYVTEASRLTVLLPLAAAERDLAPGFLEAFATAALEPGDAAAALTLLLLYEPRQ  | 553 |
| Db | 241 | VEILPVPYVTEASRLTVLLPLAAAERDLAPGFLEAFATAALEPGDAAAALTLLLLYEPRQ  | 300 |

[illegible]